



[illegible]

us-09-925-548-6.rng

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[illegible]

476	14.2	74.7	2542	21	AAX82956	Human keratinocyte
477	14.2	74.7	2712	24	AAN72976	Human H22p partial
478	14.2	74.7	2968	22	AA166822	Human protein kinase
479	14.2	74.7	2988	23	AB129735	Human P311a mediator
480	14.2	74.7	3127	21	AA651879	Arabidopsis thaliana
481	14.2	74.7	3166	23	AA864118	rat sequences thaliana
482	14.2	74.7	3419	19	AA143076	Wistar-Kyoto rat
483	14.2	74.7	3419	19	AAV34112	Human thymidine
484	14.2	74.7	3563	23	AB125225	Human thymidine
485	14.2	74.7	3429	23	AA883936	Human thymidine
486	14.2	74.7	3429	23	AA884714	Human thymidine
487	14.2	74.7	3437	17	AA106711	Human thymidine
488	14.2	74.7	3437	17	AA106711	Human thymidine
489	14.2	74.7	3437	17	AA106711	Human thymidine
490	14.2	74.7	3437	17	AA106711	Human thymidine
491	14.2	74.7	3437	17	AA106711	Human thymidine
492	14.2	74.7	3437	17	AA106711	Human thymidine
493	14.2	74.7	3437	17	AA106711	Human thymidine
494	14.2	74.7	3437	17	AA106711	Human thymidine
495	14.2	74.7	3437	17	AA106711	Human thymidine
496	14.2	74.7	3437	17	AA106711	Human thymidine
497	14.2	74.7	3437	17	AA106711	Human thymidine
498	14.2	74.7	3437	17	AA106711	Human thymidine
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500	14.2	74.7	3437	17	AA106711	Human thymidine
501	14.2	74.7	3437	17	AA106711	Human thymidine
502	14.2	74.7	3437	17	AA106711	Human thymidine
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506	14.2	74.7	3437	17	AA106711	Human thymidine
507	14.2	74.7	3437	17	AA106711	Human thymidine
508	14.2	74.7	3437	17	AA106711	Human thymidine
509	14.2	74.7	3437	17	AA106711	Human thymidine
510	14.2	74.7	3437	17	AA106711	Human thymidine
511	14.2	74.7	3437	17	AA106711	Human thymidine
512	14.2	74.7	3437	17	AA106711	Human thymidine
513	14.2	74.7	3437	17	AA106711	Human thymidine
514	14.2	74.7	3437	17	AA106711	Human thymidine
515	14.2	74.7	3437	17	AA106711	Human thymidine
516	14.2	74.7	3437	17	AA106711	Human thymidine
517	14.2	74.7	3437	17	AA106711	Human thymidine
518	14.2	74.7	3437	17	AA106711	Human thymidine
519	14.2	74.7	3437	17	AA106711	Human thymidine
520	14.2	74.7	3437	17	AA106711	Human thymidine

[illegible]











... detection an inflammation (especially chronic) in a tissue or an effector  
... response in a subject, exposure of a subject to a pathogen or several  
... infectious disease (e.g., psoriasis, rheumatoid arthritis,  
... to several of these, asthma, the asthma, certain types of allergy, tumor,  
... respiratory bowel disease, AIDS, adult respiratory distress syndrome,  
... inflammatory bowel disease, Crohn's Disease, ulcerative colitis,  
... peritoneal disease; also bacterial infection, viral infection,  
... parasitic infection, protozoan infection, fungal infection, and/or  
... action for treating one of the above conditions, the present  
... N for the sequence data for this patent did not claim part  
... of the printed specification, but was defined in electronic  
... format directly from WIP0 at  
... [http://www.who.int/genbank/published\\_pat\\_sequences](http://www.who.int/genbank/published_pat_sequences).

XX Sequence: 199 bp; 445 A; 480 G; 476 C; 6 other

Query Match: 100.0% Score 19; 19 24; 19 24; 19 24

Best Local Similarity: 100.0%; Prod. No. 14

Matches 19; Conservation 0; Mismatches 0; 1 4 2 2 0 0 0 0 0

27 1 11AAAGAGGATGAAAGG 19

14 1 1111111111111111 19

14 1 11AAAGAGGATGAAAGG 19

1591111

AAAGAGGATG

XX AAAGAGGATG standard; DNA: 1191 bp.

XX AAAGAGGATG

XX 01 N-V 2001 (first entry)

XX human immune/hematopoietic antigen genomic sequence, EP 1,191,449.

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t is about capable of modulating GCA or an inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease using the gene expression profile; (4) detecting (M4) an inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease, by detecting the level of expression in a sample of the tissue of gene(s) from GS, where (4) treating (M5) an inflammation (especially chronic) or in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease, by contacting a tissue having inflammation with an agent that modulates the expression of gene(s) from GS in the tissue, M1 is useful for detecting GCA; M2 is useful for modulating GCA; M3 is useful for detecting an inflammation in a tissue; M4 is useful for detecting an inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease (e.g., psoriasis, rheumatoid arthritis, thrombotic thrombocytopenic purpura, asthma, adult respiratory distress syndrome, hemophagocytic lymphohistiocytosis, Kawasaki's disease, viral infection, inflammatory bowel disease, Crohn's disease, ulcerative colitis, peritonitis, infectious mononucleosis, hepatitis, HIV/AIDS, bacterial infection, fungal infection, parasitic infection, protozoal infection, rickettsial infection, etc.); also for treating one of the above conditions. The present sequence represents a gene differentially expressed in granulocytes. Note: the sequence data for this patent did not form part of the claimed specification, but was obtained in electronic format directly from WHO at <http://www.who.int/genbank>.

[illegible]

RESULTS 2M	
ABU72171	
110	ABU72171 standard; cDNA; 260 bp.
XX	
XX	A
XX	ABU72171;
XX	
101	14 MAY-2002 (first entry)
XX	
XX	corn tassel-derived polynucleotide (cdps) SEQ ID NO:1545.
XX	
XX	corn: corn tassel-derived polynucleotide; cdps; hybrid breeding; CDPS;
KW	thelence; character; age; development; disease resistance;
KW	environmental adaptability; quality; yield; molecular marker;
KW	multivariate trait; plant breeding; corn tassel; gene; ss.
XX	
XX	25-34 days.
XX	
101	US2001051345-A1.
XX	
XX	1014P-2001.
101	
XX	1014P-1999; 990S-0294094.
101	
XX	21-APR-1998; 980S-082567P.
XX	
XX	(CAL-2) LALCUDI R V.
101	(11-42) LFC L Y.
101	(SHERZ) SHERAN R K.
XX	
101	Lalacudi RV, Ito LY, Sherman RK;
XX	
101	WPI; 2002-163647/21.
101	

XX Novel purified corn tassel-derived polynucleotide used at least one method for  
XX determining altered gene expression, to recover regulatory elements and  
XX to follow inheritance of desirable characteristics through genetic  
XX breeding programs

XX Claim 1; SEQ ID NOs: 2-11; English.

XX The present sequence describes a purified corn tassel derived  
XX polynucleotide sequence (cops) comprising a nucleic acid segment  
XX selected from those given in ABL70924 to ABL76834, the cops which are  
XX encodable for determining altered gene expression, the cops which can  
XX be used for determining altered gene expression, the cops which  
XX regulate genes and to follow inheritance of desirable  
XX characteristics through hybrid breeding programs. (1) are also associated  
XX with plant development, disease resistance, environmental  
XX adaptability, quality and yield, and as molecular markers for the  
XX inheritance of multiple traits in a plant breeding program. (2) are  
XX used to produce a tassel specific profile of gene transferable  
XX transmittance to clone regulatory elements for use in transgenic  
XX vectors, improve a polyphyletic identity, isolate or identify  
XX identical or related corn tassel nucleic acid sequences from DNA  
XX libraries. In nucleic acid hybridization or amplification by PCR,  
XX as query sequences to determine homology of known sequences, and  
XX for use in Southern or Northern hybridisation, and to identify the  
XX presence of and/or to determine the degree of similarity between two  
XX (or more) nucleic acid sequences.

XX Sequence: 269 bp; 72 A's; 66 C's; 69 G's; 60 T's; no other;

XX Query Match: BLAST Score: 15.4; E(-24); Length: 250;  
XX Best Local Similarity: 94.1%; Pos.: No. 1, 40-92;  
XX Matches (%): Conservation of: Bblastdb S L; Index:

XX X GAAGACGATCAAAAGA 19  
XX UUUUUUUUUU 414  
XX 125 GGGGGGGGGGGGAAG 141

XX RESULT 29  
XX ABL70924  
XX 10 ABL70924 standard cDNA: 250 bp.  
XX AC ABL70924:  
XX 1A-MAY 2002 (first entry)  
XX Corn tassel-derived polynucleotide (cops) SEQ ID NO: 299.  
XX Corn tassel-derived polynucleotide (cops): hybrid for identifying  
XX inheritance; characterized strength development; disease resistance;  
XX environmental adaptability; quality; yield; molecular marker;  
XX multigene trait; plant breeding; corn tassel; gene SS.  
XX Zea mays.  
XX US20020061466 A1.  
XX 1A-JUN 2002.  
XX PF 16-APR 1999: 99US 6204000.  
XX ZI-AUG-1999: 99US 092574P.  
XX (LAI//) LAUREN K V.  
XX (DOL//) DOLL Y.  
XX (SHER//) SHERMAN R K.  
XX Lalandi BV, Ho LY, Shortall PK  
XX WPI: 2002 143647ZL.  
XX XX









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